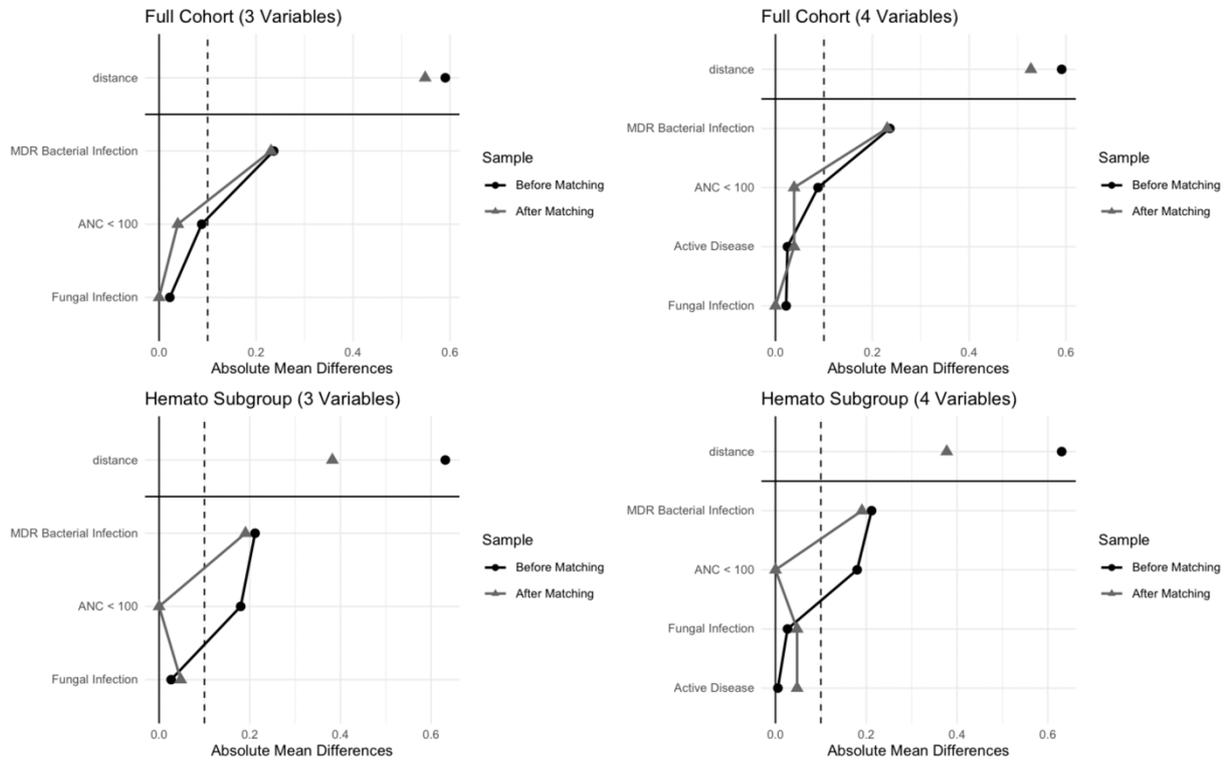


Figure S5. Covariate Balance Before and After Propensity Score Matching in Full Cohort and High-Risk Hematologic Subgroup



Love plots display standardized mean differences (SMDs) before and after 1:1 nearest-neighbor matching across four models:

- (1) Full cohort matched on 3 variables (ANC <100, fungal infection, MDR bacterial infection);
- (2) Full cohort matched on 4 variables (adds active disease status);
- (3) High-risk hematologic subgroup matched on 3 variables;
- (4) High-risk subgroup matched on 4 variables.

The dashed vertical line indicates the threshold for acceptable balance (SMD < 0.1).

All models demonstrated an improved post-matching covariate balance, supporting the robustness of PSM-adjusted analyses.